CORRECTED SEQUENCE LISTING

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<120> METHODS OF INHIBITING T CELL PROLIFERATION OR IL-2 ACCUMULATION WITH CTLA-4 SPECIFIC ANTIBODIES (AS AMENDED)

<130> RPI-016CPA2DV

<140> 10/732847
<141> 2003-12-09

<150> 08/253783
<151> 1994-06-03

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Thr Val His Thr
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Val Pro His Ala
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Gly Ala Phe Arg
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<223> Synthetic peptide

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Asp Gln Pro Phe
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Val Leu Thr Phe Leu Glu Arg Leu Leu Pro Ala Val Val Pro Arg Ser
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Cys His Pro Gly
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<210> 27
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<400> 27
Leu Ser Trp Gly Leu Glu Pro Trp Glu Gly Ser Phe Leu Trp Leu Thr
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Glu Ser Pro Met
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<210> 28
<211> 20
<212> PRT
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Pro Asp Ser Cys
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<210> 29
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<220>
<223> Oligonucleotide

<400> 31
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38, 39, 40, 41, 42, 43, 44, 45, 46
<223> Xaa = Any Amino Acid

<223> Synthetic peptide

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37, 38, 39, 40, 41, 42, 43, 44, 45
<223> Xaa = Any Amino Acid

<223> Synthetic peptide

<400> 34
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20 25 30
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35 40 45

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atc gcc agc ttt gtg tgt cag tat gca tct cca ggc aaa gcc act gag 96
Ile Ala Ser Phe Val Cys Gln Tyr Ala Ser Pro Gly Lys Ala Thr Glu
20 25 30

gtc cgg gtg aca gtg ctt cgg cag gct gac agc cag gtg act gaa gtc 144
Val Arg Val Thr Val Leu Arg Gln Ala Asp Ser Gln Val Thr Glu Val
35 40 45

tgt gcg gca acc tac atg atg ggg aat gag ttg acc ttc cta gat gat 192
Cys Ala Ala Thr Tyr Met Met Gly Asn Glu Leu Thr Phe Leu Asp Asp
50 55 60

tcc atc tgc acg ggc acc tcc agt gga aat caa gtg aac ctc act atc 240
Ser Ile Cys Thr Gly Thr Ser Ser Gly Asn Gln Val Asn Leu Thr Ile
65 70 75 80

caa gga ctg agg gcc atg gac acg gga ctc tac atc tgc aag gtg gag 288
Gln Gly Leu Arg Ala Met Asp Thr Gly Leu Tyr Ile Cys Lys Val Glu
85 90 95

ctc atg tac cca ccg cca tac tac ctg ggc ata ggc aac gga acc cag 336
Leu Met Tyr Pro Pro Pro Tyr Tyr Leu Gly Ile Gly Asn Gly Thr Gln
100 105 110

att tat gta att gat cca gaa ccg tgc cca gat tct gac ttc ctc ctc 384
Ile Tyr Val Ile Asp Pro Glu Pro Cys Pro Asp Ser Asp Phe Leu Leu
115 120 125

tgg atc ctt gca gca gtt agt tcg ggg ttg ttt ttt tat agc ttt ctc 432
Trp Ile Leu Ala Ala Val Ser Ser Gly Leu Phe Phe Tyr Ser Phe Leu
130 135 140

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ctc aca gct gtt tct ttg agc aaa atg cta aag aaa aga agc cct ctt   480
Leu Thr Ala Val Ser Leu Ser Lys Met Leu Lys Lys Arg Ser Pro Leu
145                 150                 155                 160

aca aca ggg gtc tat gtg aaa atg ccc cca aca gag cca gaa tgt gaa   528
Thr Thr Gly Val Tyr Val Lys Met Pro Pro Thr Glu Pro Glu Cys Glu
                165                 170                 175

aag caa ttt cag cct tat ttt att ccc atc aat                      561
Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn
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<210> 36
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<220>
<223> Protein

<400> 36

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        35                  40                  45
Cys Ala Ala Thr Tyr Met Met Gly Asn Glu Leu Thr Phe Leu Asp Asp
    50                  55                  60
Ser Ile Cys Thr Gly Thr Ser Ser Gly Asn Gln Val Asn Leu Thr Ile
65                  70                  75                  80
Gln Gly Leu Arg Ala Met Asp Thr Gly Leu Tyr Ile Cys Lys Val Glu
                85                  90                  95
Leu Met Tyr Pro Pro Pro Tyr Tyr Leu Gly Ile Gly Asn Gly Thr Gln
            100                 105                 110
Ile Tyr Val Ile Asp Pro Glu Pro Cys Pro Asp Ser Asp Phe Leu Leu
        115                 120                 125
Trp Ile Leu Ala Ala Val Ser Ser Gly Leu Phe Phe Tyr Ser Phe Leu
    130                 135                 140
Leu Thr Ala Val Ser Leu Ser Lys Met Leu Lys Lys Arg Ser Pro Leu
145                 150                 155                 160
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Lys Gln Phe Gln Pro Tyr Phe Ile Pro Ile Asn
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ccaaccagcg atggccgcag caatgcacgt ggcccagcct gctgtgg 107